## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/588,052
Source:	IFWP,
Date Processed by STIC:	8/10/06

## ENTERED



**IFWP** 

RAW SEQUENCE LISTING DATE: 08/10/2006
PATENT APPLICATION: US/10/588,052 TIME: 11:05:24

Input Set: A:\44352-0011-00-US sequence listing(pct 05 235).txt

Output Set: N:\CRF4\08102006\J588052.raw

```
3 <110> APPLICANT: Lifenza Co., Ltd.
     5 <120> TITLE OF INVENTION: PROTEIN WITH ACTIVITY OF HYDROLYZING AMYLOPECTIN, STARCH,
             GLYCOGEN AND AMYLOSE, GENE ENCODING THE SAME, CELL EXPRESSING THE
             SAME, AND PRODUCTION METHOD THEREOF
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/588,052
C--> 9 <141> CURRENT FILING DATE: 2006-07-31
W--> 0 <130> FILE REFERENCE:
     9 <150> PRIOR APPLICATION NUMBER: KR2004-0006186
    10 <151> PRIOR FILING DATE: 2004-01-30
    12 <160> NUMBER OF SEQ ID NOS: 4
    14 <170> SOFTWARE: KopatentIn 1.71
    16 <210> SEQ ID NO: 1
    17 <211> LENGTH: 647
    18 <212> TYPE: PRT
    19 <213> ORGANISM: Artificial Sequence
    21 <220> FEATURE:
    22 <223> OTHER INFORMATION: E. coli BL21(DE3)pLysS
    25 <400> SEQUENCE: 1
    26 Met Leu Ile Asn Phe Phe Ile Ala Val Leu Gly Val Ile Ser Leu
    27 1
                         5
                                             10
    29 Ser Pro Ile Val Val Ala Arg Tyr Ile Leu Arg Arg Asp Cys Thr Thr
                                         25
    32 Val Thr Val Leu Ser Ser Pro Glu Ser Val Thr Ser Ser Asn His Val
                35
                                     40
    35 Glu Leu Ala Ser His Glu Met Cys Asp Ser Thr Leu Ser Ala Ser Leu
    38 Tyr Ile Tyr Asn Asp Asp Tyr Asp Lys Ile Val Thr Leu Tyr Tyr Leu
                            70
                                                 75
    41 Thr Ser Ser Gly Thr Thr Gly Ser Val Thr Ala Ser Tyr Ser Ser Ser
    44 Leu Ser Asn Asn Trp Glu Leu Trp Ser Leu Ser Ala Pro Ala Ala Asp
                   100
                                        105
    47 Ala Val Glu Ile Thr Gly Ala Ser Tyr Val Asp Ser Asp Ala Ser Ala
                                   120
               115
    50 Thr Tyr Ala Thr Ser Phe Asp Ile Pro Leu Thr Thr Thr Thr Ser
                               135
    53 Ser Ser Ser Ala Ser Ala Thr Ser Thr Ser Ser Leu Thr Thr Thr Ser
                                                155
                           150
    56 Ser Val Ser Ile Ser Val Ser Val Pro Thr Gly Thr Ala Ala Asn Trp
                                            170
    59 Arg Gly Arg Ala Ile Tyr Glu Ile Val Thr Asp Arg Phe Ala Arg Thr
                   180
                                        185
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62 Asp Gly Ser Thr Thr Tyr Leu Cys Asp Val Thr Asp Arg Val Tyr Cys

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63 195 200 65 Gly Gly Ser Tyr Glu Gly Ile Ile Asn Met Leu Asp Tyr Ile Glu Gly 215 220 68 Met Gly Phe Thr Ala Ile Trp Ile Ser Pro Ile Val Glu Asn Ile Pro 230 235 71 Asp Asp Thr Gly Tyr Gly Tyr Ala Tyr His Gly Tyr Trp Met Lys Asp 74 Ile Phe Ala Leu Asn Thr Asn Phe Gly Thr Ala Asp Asp Leu Ile Ala 260 265 77 Leu Ala Thr Glu Leu His Asn Arg Gly Met Tyr Leu Met Val Asp Ile 280 80 Val Val Asn His Phe Ala Phe Ser Gly Ser His Ala Asp Val Asp Tyr 295 83 Ser Glu Tyr Phe Pro Tyr Ser Ser Glu Asp Tyr Phe His Ser Phe Cys 310 315 86 Trp Ile Thr Asp Tyr Ser Asn Glu Thr Asn Val Glu Gln Cys Trp Leu 325 330 89 Gly Asp Asp Thr Val Pro Leu Val Asp Val Asn Thr Glu Leu Asp Thr 340 345 92 Val Lys Ser Glu Tyr Gln Ser Trp Val Glu Glu Leu Ile Ala Asn Tyr 360 95 Ser Ile Asp Gly Leu Arg Ile Asp Thr Val Lys His Val Glu Met Asp 375 98 Phe Trp Ala Pro Phe Glu Glu Ala Ala Gly Ile Tyr Ala Val Gly Glu 390 101 Val Phe Asp Gly Asp Pro Ser Tyr Thr Cys Pro Tyr Glu Glu Asn Leu 405 410 104 Asp Gly Val Leu Asn Tyr Pro Val Tyr Tyr Pro Val Val Ser Ala Phe 425 107 Glu Ser Val Ser Gly Ser Val Ser Ser Leu Val Asp Met Ile Asp Thr 435 440 110 Leu Lys Ser Glu Cys Thr Asp Thr Thr Leu Leu Gly Ser Phe Leu Glu 455 113 Asn Glu Asp Asn Pro Arg Phe Pro Ser Tyr Thr Ser Asp Glu Ser Leu 470 475 116 Ile Lys Asn Ala Ile Ala Phe Thr Met Leu Ser Asp Gly Ile Pro Ile 117 485 490 119 Ile Tyr Tyr Gly Glu Glu Gln Gly Leu Asn Gly Gly Asn Asp Pro Tyr 505 500 122 Asn Arg Glu Ala Leu Trp Leu Thr Gly Tyr Ser Thr Thr Ser Thr Phe 515 520 125 Tyr Lys Tyr Ile Ala Ser Leu Asn Glu Ile Arg Asn Glu Ala Ile Tyr 535 128 Lys Asp Asp Thr Tyr Leu Thr Tyr Glu Asn Trp Val Ile Tyr Ser Asp 129 545 550 555 131 Ser Thr Thr Ile Ala Met Arg Lys Gly Phe Thr Gly Asn Glu Ile Ile 570 134 Thr Val Leu Ser Asn Leu Gly Thr Ser Gly Ser Ser Tyr Thr Leu Thr 580 585

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Output Set: N:\CRF4\08102006\J588052.raw

```
137 Leu Ser Asn Thr Gly Tyr Thr Ala Ser Ser Val Val Tyr Glu Ile Leu
                                600
            595
                                                     605
140 Thr Cys Thr Ala Val Thr Val Asp Ser Ser Gly Asn Leu Ala Val Pro
        610
                            615
141
143 Met Ser Ser Gly Leu Pro Lys Val Phe Tyr Glu Glu Ser Gln Leu Val
                                             635
146 Gly Ser Gly Ile Cys Ser Met
147
150 <210> SEQ ID NO: 2
151 <211> LENGTH: 1946
152 <212> TYPE: DNA
153 <213> ORGANISM: Artificial Sequence
155 <220> FEATURE:
156 <223 > OTHER INFORMATION: E. coli BL21(DE3)pLysS
159 <400> SEQUENCE: 2
160 atgttgctga tcaacttttt catcgctgtt ctgggagtga tatcactgtc tcctattgtg
                                                                               60
162 gttgctcgtt atattcttcg acgagattgc actacagtta cggtcttgtc ctcccctgag
                                                                              120
164 tctgtgacga gttcgaacca tgttcagcta gccagtcatg agatgtgcga cagtaccttg
                                                                              180
166 traggegtere tttatateta caatgatgat tatgataaga ttgtgacact ttattatett
                                                                              240
                                                                              300
168 acategtegg geacaactgg gteegtaaca gegtettatt ettetagttt gagtaacaac
170 tgggaattgt ggtctctctc ggctccggct gcagatgctg tcgagatcac tggagctagt
                                                                              360
172 tatgtagaca gcgatgcatc tgcgacatac gccacgtctt ttgatatacc tcttactacc
                                                                              420
174 acgacaacgt cgtcgtcttc tgctagtgcg acttcaacat ctagtctaac cacaacatct
                                                                              480
176 aqtqtttcca tttcqqtqtc cqtccctaca qqaacaqctq caaattqqcq aggtagggct
                                                                              540
178 atctatcaqa tcqtqactqa taqatttqca cqcactqacq qctccaccac atatttatqc
                                                                              600
180 gatgttaccg atagggtcta ttgcggaggg tcttatcagg ggattatcaa tatgctggat
                                                                              660
182 tacatccaag gcatgggctt tactgctatt tggatttctc ctatagtgga aaatattccc
                                                                              720
184 gatgacaccg gatacggtta cgcatatcat ggttattgga tgaaagatat cttcgccctg
                                                                              780
186 aatacaaatt ttggtactgc agacgatttg atagcgttgg ctacggaatt gcataatcgc
                                                                              840
188 ggcatgtact tgatggttga tattgttgtc aatcactttg ctttctcagg aagtcatgcc
                                                                              900
190 gacgtggact actctgaata tttcccgtat tcgtcccagg attattttca ttcattttgc
                                                                              960
192 tggattacag attactcgaa tcagacaaac gttgagcagt gctggcttgg cgacgatact
                                                                             1020
194 gttcctctcg tggacgtcaa tacccaactt gacaccgtga aaagtgaata tcaatcctgg
                                                                             1080
196 gttcaagaac ttatagctaa ttactctatt gacggcctaa gaattgacac cgtcaagcac
                                                                             1140
198 gtgcagatgg atttttgggc accatttcaa gaggctgcag ggatttacgc cgttggtgaa
                                                                             1200
200 gtattcgacg gtgatccatc ctacacatgt ccatatcagg aaaatcttga cggtgtcttg
                                                                             1260
202 aattatcctg tttattatcc tgtcgtctct gcgtttgaga gtgttagtgg gtcggtctcc
                                                                             1320
204 tegttagteg atatgattga taegeteaag tetgaatgea eegacaetae teteetagge
                                                                             1380
206 teetttetag agaateaaga taateegega tteeetaget acaettetga tgagtettta
                                                                             1440
208 attaaaaatg cgatcgcttt cactatgctc tcagacggca ttcccataat ttattacggt
                                                                             1500
210 caggagcaag gcctcaatgg tggaaacgat ccctataatc gagaggcgct ttggcttacg
                                                                             1560
212 ggctactcca caacgtcgac gttctacaaa tacattgcgt cgttgaatca gattagaaat
                                                                             1620
214 caggetatat acaaagatga tacttatete acatateaga actgggttat ttatteggat
                                                                             1680
216 tccacgacaa tagcaatgcg gaaaggtttt acagggaacc aaataattac ggttctgtca
                                                                             1740
218 aatcttggga ccagtggcag ttcgtacact ttgacgcttt cgaatacggg atataccgca
                                                                             1800
                                                                             1860
220 tctaqcqttq tatatqagat cttqacatqc acaqctqtga ctgtggattc gtctgggaat
222 ttqqcaqtqc cqatqtccaq tqqcctacca aaaqtctttt atcaqqaatc gcaactgqtt
                                                                             1920
                                                                             1946
224 ggctctggaa tctgctccat gtagag
227 <210> SEQ ID NO: 3
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RAW SEQUENCE LISTING

DATE: 08/10/2006

PATENT APPLICATION: US/10/588,052

TIME: 11:05:24

Input Set : A:\44352-0011-00-US sequence listing(pct 05 235).txt

Output Set: N:\CRF4\08102006\J588052.raw

- 228 <211> LENGTH: 27
- 229 <212> TYPE: DNA
- 230 <213> ORGANISM: Artificial Sequence
- 232 <220> FEATURE:
- 233 <223> OTHER INFORMATION: L. starkeyi primer 1(sense)
- 236 <400> SEQUENCE: 3
- 237 tacagttacg gtcttgtcct cccctga

27

- 240 <210> SEQ ID NO: 4
- 241 <211> LENGTH: 21
- 242 <212> TYPE: DNA
- 243 <213> ORGANISM: Artificial Sequence
- 245 <220> FEATURE:
- 246 <223> OTHER INFORMATION: L. starkeyi primer 2(antisense)
- 249 <400> SEQUENCE: 4
- 250 ctctacatgg agcagattcc a

21

VERIFICATION SUMMARY

DATE: 08/10/2006

PATENT APPLICATION: US/10/588,052

TIME: 11:05:25

Input Set : A:\44352-0011-00-US sequence listing(pct 05 235).txt

Output Set: N:\CRF4\08102006\J588052.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE